

#398943 v1 - A34584-A-PCT-USA Sequence Listing SEQUENCE LISTING

<110> Fisher, Paul B. Leszcyniecka, Magdalena

<120> GENES DISPLAYING ENHANCED EXPRESSION DURING CELLULAR SENESCENCE AND TERMINAL CELL DIFFERENTIATION AND USES THEREOF

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<130> A34584-A-PCT-USA (070050.1664)
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<141> 2000-02-02
<150> US 09/243,277
<151> 1999-02-02
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#398943 v1 - A34584-A-PCT-USA Sequence Listing
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<222> 434, 459, 460
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<213> Homo sapiens
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#398943 v1 - A34584-A-PCT-USA Sequence Listing
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#398943 v1 - A34584-A-PCT-USA Sequence Listing
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<221> unsure
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#398943 v1 - A34584-A-PCT-USA Sequence Listing
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Phe Ala Asp Gly Ser Ala Val Val Gln Ser Gly Asp Thr Ala Val Met 50 60 Val Thr Ala Val Ser Lys Thr Lys Pro Ser Pro Ser Gln Phe Met Pro 65 70 75 80
Leu Val Val Asp Tyr Arg Gln Lys Ala Ala Ala Ala Gly Arg Ile Pro
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#398943 v1 - A34584-A-PCT-USA Sequence_Listing Thr Asn Tyr Leu Arg Arg Glu Val Gly Thr Ser Asp Lys Glu Ile Leu 100 105 110 Thr Ser Arg Ile Ile Asp Arg Ser Ile Arg Pro Leu Phe Pro Ala Gly 115 120 125 Tyr Phe Tyr Asp Thr Gln Val Leu Cys Asn Leu Leu Ala Val Asp Gly
130 135 140 Val Asn Glu Pro Asp Val Leu Ala Ile Asn Gly Ala Ser Val Ala Leu 145 150 155 160 Ser Leu Ser Asp Ile Pro Trp Asn Gly Pro Val Gly Ala Val Arg Ile 165 170 Gly Ile Ile Asp Gly Glu Tyr Val Val Asn Pro Thr Arg Lys Glu Met 180 185 Ser Ser Ser Thr Leu Asn Leu Val Val Ala Gly Ala Pro Lys Ser Gln 195 200 205 Ile Val Met Leu Glu Ala Ser Ala Glu Asn Ile Leu Gln Gln Asp Phe 215 Cys His Ala Ile Lys Val Gly Val Lys Tyr Thr Gln Gln Ile Ile Gln 235 230 Gly Ile Gln Gln Leu Val Lys Glu Thr Gly Val Thr Lys Arg Thr Pro 250 245 Gln Lys Leu Phe Thr Pro Ser Pro Glu Ile Val Lys Tyr Thr His Lys 260 270 265 Leu Ala Met Glu Arg Leu Tyr Ala Val Phe Thr Asp Tyr Glu His Asp 275 280 285 Lys Val Ser Arg Asp Glu Ala Val Asn Lys Ile Arg Leu Asp Thr Glu 290 295 300 Glu Gln Leu Lys Glu Lys Phe Pro Glu Ala Asp Pro Tyr Glu Ile Ile 305 310 315 320 Glu Ser Phe Asn Val Val Ala Lys Glu Val Phe Arg Ser Ile Val Leu 325 330 Asn Glu Tyr Lys Arg Cys Asp Gly Arg Asp Leu Thr Ser Leu Arg Asn 345 Val Ser Cys Glu Val Asp Met Phe Lys Thr Leu His Gly Ser Ala Leu 355 360 365 360 Phe Gln Arg Gly Gln Thr Gln Val Leu Cys Thr Val Thr Phe Asp Ser 370 375 Leu Glu Ser Gly Ile Lys Ser Asp Gln Val Ile Thr Ala Ile Asn Gly 390 395 400 Ile Lys Asp Lys Asn Phe Met Leu His Tyr Glu Phe Pro Pro Tyr Ala 405 410 Thr Asn Glu Ile Gly Lys Val Thr Gly Leu Asn Arg Arg Glu Leu Gly
420 425 430 His Gly Ala Leu Ala Glu Lys Ala Leu Tyr Pro Val Ile Pro Arg Asp
435 440 445 Phe Pro Phe Thr Ile Arg Val Thr Ser Glu Val Leu Glu Ser Asn Gly 455 460 Ser Ser Ser Met Ala Ser Ala Cys Gly Gly Ser Leu Ala Leu Met Asp 470 475 Ser Gly Val Pro Ile Ser Ser Ala Val Ala Gly Val Ala Ile Gly Leu 485 490 Val Thr Lys Thr Asp Pro Glu Lys Gly Glu Ile Glu Asp Tyr Arg Leu
500 505 510 Leu Thr Asp Ile Leu Gly Ile Glu Asp Tyr Asn Gly Asp Met Asp Phe
515 520 525 Lys Ile Ala Gly Thr Asn Lys Gly Ile Thr Ala Leu Gln Ala Asp Ile 535 Lys Leu Pro Gly Ile Pro Ile Lys Ile Val Met Glu Ala Ile Gln Gln 550 555 Ala Ser Val Ala Lys Lys Glu Ile Leu Gln Ile Met Asn Lys Thr Ile 565 570 575 Ser Lys Pro Arg Ala Ser Arg Lys Glu Asn Gly Pro Val Val Glu Thr 58Ŏ 585 590

#398943 v1 - A34584-A-PCT-USA Sequence Listing Val Gln Val Pro Leu Ser Lys Arg Ala Lys Phe Val Gly Pro Gly Gly Tyr Asn Leu Lys Lys Leu Gln Ala Glu Thr Gly Val Thr Ile Ser Gln Val Asp Glu Glu Thr Phe Ser Val Phe Ala Pro Thr Pro Ser Val Met His Glu Ala Arg Asp Phe Ile Thr Glu Ile Cys Lys Asp Asp Gln Glu Gln Gln Leu Glu Phe Gly Ala Val Tyr Thr Ala Thr Ile Thr Glu Ile Arg Asp Thr Gly Val Met Val Lys Leu Tyr Pro Asn Met Thr Ala Val Leu Leu His Asn Thr Gln Leu Asp Asn Glu Arg Leu Asn Ile Leu Leu Pro

<210> 43 <211> 705 <212> PRT

<213> Bacillus subtilis

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#398943 v1 - A34584-A-PCT-USA Sequence Listing
                                              315
Gly Val Asp Gln Ile Arg Pro Leu Ser Ser Glu Val Gly Leu Leu Pro
                  325
                                         330
                                                                335
Arg Thr His Gly Ser Gly Leu Phe Thr Arg Gly Gln Thr Gln Ala Leu 340 345 350
Ser Val Cys Thr Leu Gly Ala Leu Gly Asp Val Gln Ile Leu Asp Gly 355 _ _ 360 . . . . 365
Leu Gly Val Glu Glu Ser Lys Arg Phe Met His His Tyr Asn Phe Pro 370 380
Gln Phe Ser Val Gly Glu Thr Gly Pro Met Arg Gly Pro Gly Arg Arg
                      390
                                             395
Glu Ile Gly His Gly Ala Leu Gly Glu Arg Ala Leu Glu Pro Val Ile
405 410 415
Pro Ser Glu Lys Asp Phe Pro Tyr Thr Val Arg Leu Val Ser Glu Val 420 425 430
Leu Glu Ser Asn Gly Ser Thr Ser Gln Ala Ser Ile Cys Ala Ser Thr
435 440 445
Leu Ala Met Met Asp Ala Gly Val Pro Ile Lys Ala Pro Val Ala Gly
                           455
Ile Ala Met Gly Leu Val Lys Ser Gly Glu His Tyr Thr Val Leu Thr
                      470
                                             475
Asp Ile Gln Gly Met Glu Asp Ala Leu Gly Asp Met Asp Phe Lys Val
485 490 495
Ala Gly Thr Glu Lys Gly Val Thr Ala Leu Gln Met Asp Ile Lys Ile 500 505 510
Glu Gly Leu Ser Arg Glu Ile Leu Glu Glu Ala Leu Gln Gln Ala Lys
                                520
         515
Lys Gly Arg Met Glu Ile Leu Asn Ser Met Leu Ala Thr Leu Ser Glu
    530
                           535
                                                  540
Ser Arg Lys Glu Leu Ser Arg Tyr Ala Pro Lys Ile Leu Thr Met Thr 545 550 550 560
ile Asn Pro Asp Lys ile Arg Asp Val Ile Gly Pro Ser Gly Lys Gln
565 575
Ile Asn Lys Ile Ile Glu Glu Thr Gly Val Lys Ile Asp Ile Glu Gln
580 585 590
Asp Gly Thr Ile Phe Ile Ser Ser Thr Asp Glu Ser Gly Asm Glm Lys 595 600 605
Ala Lys Lys Ile Ile Glu Asp Leu Val Arg Glu Val Glu Val Gly Gln
    610
                           615
                                                  620
Leu Tyr Leu Gly Lys Val Lys Arg Ile Glu Lys Phe Gly Ala Phe Val 625 630 635 640
Glu Ile Phe Ser Gly Lys Asp Gly Leu Val His Ile Ser Glu Leu Ala
645 650 655
Leu Glu Arg Val Gly Lys Val Glu Asp Val Val Lys Ile Gly Asp Glu
660 665 670
Ile Leu Val Lys Val Thr Glu Ile Asp Lys Gln Gly Arg Val Asn Leu 675 _ 680 685
Ser Arg Lys Ala Val Leu Arg Glu Glu Lys Glu Lys Glu Glu Gln Gln 690 695 700
Ser
705
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<210> 44 <211> 705 <212> PRT <213> Homo sapiens

#398943 v1 - A34584-A-PCT-USA Sequence_Listing Asp Leu Gly Asn Arg Lys Leu Glu Ile Ser Ser Gly Lys Leu Ala Arg 40 45 Phe Ala Asp Gly Ser Ala Val Val Gln Ser Gly Asp Thr Ala Val Met 50 55 60 Val Thr Ala Val Ser Lys Thr Lys Pro Ser Pro Ser Gln Phe Met Pro 65 70 75 80 Leu Val Val Asp Tyr Arg Gln Lys Ala Ala Ala Ala Gly Arg Ile Pro 85 90 95 Thr Asn Tyr Leu Arg Arg Glu Val Gly Thr Ser Asp Lys Glu Ile Leu 105 100 Thr Ser Arg Ile Ile Asp Arg Ser Ile Arg Pro Leu Phe Pro Ala Gly 115 120 Tyr Phe Tyr Asp Thr Gln Val Leu Cys Asn Leu Leu Ala Val Asp Gly 135 140 Val Asn Glu Pro Asp Val Leu Ala Ile Asn Gly Ala Ser Val Ala Leu 145 150 155 160 Ser Leu Ser Asp Ile Pro Trp Asn Gly Pro Val Gly Ala Val Arg Ile 165 170 175 Gly Ile Ile Asp Gly Glu Tyr Val Val Asn Pro Thr Arg Lys Glu Met 185 Ser Ser Ser Thr Leu Asn Leu Val Val Ala Gly Ala Pro Lys Ser Gln 205 195 200 Ile Val Met Leu Glu Ala Ser Ala Glu Asn Ile Leu Gln Gln Asp Phe 210 215 220 Cys His Ala Ile Lys Val Gly Val Lys Tyr Thr Gln Gln Ile Ile Gln 225 230 235 240 Gly Ile Gln Gln Leu Val Lys Glu Thr Gly Val Thr Lys Arg Thr Pro 245 250 255 Gln Lys Leu Phe Thr Pro Ser Pro Glu Ile Val Lys Tyr Thr His Lys 260 265 Leu Ala Met Glu Arg Leu Tyr Ala Val Phe Thr Asp Tyr Glu His Asp 275 280 285 Lys Val Ser Arg Asp Glu Ala Val Asn Lys Ile Arg Leu Asp Thr Glu 295 290 Glu Gln Leu Lys Glu Lys Phe Pro Glu Ala Asp Pro Tyr Glu Ile Ile 305 310 320 Glu Ser Phe Asn Val Val Ala Lys Glu Val Phe Arg Ser Ile Val Leu 325 330 Asn Glu Tyr Lys Arg Cys Asp Gly Arg Asp Leu Thr Ser Leu Arg Asn 340 345 350 Val Ser Cys Glu Val Asp Met Phe Lys Thr Leu His Gly Ser Ala Leu 355 360 365 Phe Gln Arg Gly Gln Thr Gln Val Leu Cys Thr Val Thr Phe Asp Ser 375 380 Leu Glu Ser Gly Ile Lys Ser Asp Gln Val Ile Thr Ala Ile Asn Gly 390 395 400 Ile Lys Asp Lys Asn Phe Met Leu His Tyr Glu Phe Pro Pro Tyr Ala 410 405 Thr Asn Glu Ile Gly Lys Val Thr Gly Leu Asn Arg Arg Glu Leu Gly 425 430 His Gly Ala Leu Ala Glu Lys Ala Leu Tyr Pro Val Ile Pro Arg Asp 440 445 Phe Pro Phe Thr Ile Arg Val Thr Ser Glu Val Leu Glu Ser Asn Gly 460 450 455 Ser Ser Ser Met Ala Ser Ala Cys Gly Gly Ser Leu Ala Leu Met Asp 470 475 480 Ser Gly Val Pro Ile Ser Ser Ala Val Ala Gly Val Ala Ile Gly Leu 485 490 495 Val Thr Lys Thr Asp Pro Glu Lys Gly Glu Ile Glu Asp Tyr Arg Leu 500 _ _ _ 510 Leu Thr Asp Ile Leu Gly Ile Glu Asp Tyr Asn Gly Asp Met Asp Phe 515 520 525 Lys Ile Ala Gly Thr Asn Lys Gly Ile Thr Ala Leu Gln Ala Asp Ile Page 18

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#398943 v1 - A34584-A-PCT-USA Sequence Listing
                         535
                                              540
Lys Leu Pro Gly Ile Pro Ile Lys Ile Val Met Glu Ala Ile Gln Gln
545
                    550
                                          555
Ala Ser Val Ala Lys Lys Glu Ile Leu Gln Ile Met Asn Lys Thr Ile
565 570 575
Ser Lys Pro Arg Ala Ser Arg Lys Glu Asn Gly Pro Val Val Glu Thr
            58Ō
                                 585
                                                      590
Val Gln Val Pro Leu Ser Lys Arg Ala Lys Phe Val Gly Pro Gly Gly
        595
                             60Ō
                                                  605
Tyr Asn Leu Lys Lys Leu Gln Ala Glu Thr Gly Val Thr Ile Ser Gln
                         615
                                              620
Val Asp Glu Glu Thr Phe Ser Val Phe Ala Pro Thr Pro Ser Val Met
                    630
                                          635
His Glu Ala Arg Asp Phe Ile Thr Glu Ile Cys Lys Asp Asp Gln Glu
                645
                                     650
Gln Gln Leu Glu Phe Gly Ala Val Tyr Thr Ala Thr Ile Thr Glu Ile
                                 665
            660
                                                      670
Arg Asp Thr Gly Val Met Val Lys Leu Tyr Pro Asn Met Thr Ala Val
        675
                             680
                                                  685
Leu Leu His Asn Thr Gln Leu Asp Asn Glu Arg Leu Asn Ile Leu Leu
                         695
                                              700
Pro
705
<210> 45
<211> 245
<212> PRT
<213> Artificial Sequence
<220>
<223> Consensus sequence between Homo sapiens OLD-35 and
      Bacillus subtilis PNPase
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Pro Phe Pro Leu Val Tyr Ala Gly Ile Pro Arg Glu Ser Lys Leu Ser
            20
Arg Ile Asp Arg Ile Arg Pro Leu Phe Gly Gln Val Val Asp Ala Gly
        35
                             40
                                                  45
Ser Ala Leu Ser Ser Asp Ile Gly Pro Val Gly Ile Asp Asn Pro Thr
50 60
                                              60
Ser Asn Leu Val Val Ala Gly Lys Ile Met Glu Ala Ala Ile Gly 65 70 75 80
Ile Val Gly Lys Lys Leu Phe Glu Leu Ala Glu Leu Glu Lys Glu Val
Glu Val Arg Ile Glu Arg Asp Gly Arg Arg Ser Glu Val His Gly Ser
            100
                                 105
                                                      110
```

Leu Phe Arg Gly Gln Thr Gln Leu Thr Leu Asp Lys Phe Met His Tyr 115 120 125 Phe Pro Glu Gly Gly Arg Arg Glu Gly His Gly Ala Leu Glu Ala Leu 135 140 Pro Val Ile Pro Asp Phe Pro Thr Arg Ser Glu Val Leu Glu Ser Asn 155 150 160 Gly Ser Ser Ala Ser Cys Leu Ala Met Asp Gly Val Pro Ile Val Ala 170 Gly Ala Gly Leu Val Glu Tyr Leu Thr Asp Ile Gly Glu Asp Gly Asp 180 185 190 Met Asp Phe Lys Ala Gly Thr Lys Gly Thr Ala Leu Gln Asp Ile Lys 195 _ _ _ 200 _ 205

Gly Ile Glu Ala Gln Gln Ala Glu Ile Leu Met Thr Ser Arg Pro Thr

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#398943 v1 - A34584-A-PCT-USA Sequence Listing Lys Gly Pro Gly Lys Glu Thr Gly Val Ile Thr Ser Ala Ile Gln Leu													
Lys 225	Gly	Pro	Gly	Lys	G1 23	u Th	r Gly	Val I	le Thr 235	Ser Ala	Ile	e Gln Leu 240	
	Val	Lys	Leu	G1u 245	23				233			240	
<210> 46 <211> 47													
<212> RNA <213> Homo sapiens													
<400> 46 uaauauuuau auauuuauau uuuuaaaaua uuuauuuauu uauuuaa													47
<210> 47 <211> 11 <212> RNA <213> Homo sapiens													
<400> 47 uauuuauuua a													11
<211 <212)> 48 L> 33 !> RN B> Ho	iA	sapie	ens									
)> 48 Iuauu		aauaı	uuua	aa	นนนน	auauuu	aau					33
<211 <212)> 49 L> 62 !> RN !> Ho	! IA	sapie	ens									
)> 49 Iuuaa		Jauul	uauu	aa	gaug	gauucu	caga	uauuua	uauuuuu	uau	uuuauuuuuu	60 62
<211 <212)> 50 L> 11 !> RN !> Ho	.1 IA	sapie	ens									
auuu)> 50 Iacau Iauca	igu g	gccau aaaua	มนนนเ ลนนนเ	uu au	uaauı uauu:	ıcgagu aaaagu	aacc aagu	cauauu cauuua	uguuuaai uacaucui	ug Jug	uauuuacauu a	60 111
<211 <212)> 51 .> 34 !> RN !> HO	A	sapie	ens									
)> 51 Iaauu		ıuauı	Juaui	uu	auuaı	Juuauu	uauu					

34

. #398943 vi - A34584-A-PCT-USA Sequence Listing